### For Debate

### **Diabetes: mellitus or lipidus?**

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### Introduction

Since ancient times the search for causes of diabetes was related to the sweetness of urine and other body fluids. In England, Thomas Willis (1621–1675) was among the first to taste the urine of diabetic patients and declare that "its sweet taste was imbued with honey sugar" with the supposition that it was derived from blood. This finding led to the addition of "mellitus" to the word diabetes.

We should reconsider whether the definition of diabetes as mellitus is justified. The traditional emphasis on the insulin-glucose axis with respect to examining glucose tolerance, although diagnostically useful does not explain the basic pathophysiological mechanisms operating in diabetes. The glucose-insulin axis has been overemphasised, while alterations in the insulin to non-esterified fatty acid (NEFA) ratio and metabolism received much less attention both as diabetes derangement and as diagnostic potential. Perhaps this is partly due to the fact that the techniques for glucose measurements have been considerably simplified for home use, whereas the measurement of NEFA and triglycerides is still encumbered with relative difficulties, being more expensive and time consuming.

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*Abbreviations:* DAG, diacylglycerol; LCFA CoA, long chain fatty acid coenzyme A; PKB, protein kinase B; PKC, protein kinase C; TZD, thiazolidinediones.

## NEFA as the main fat energy supplier, affected by diabetes and obesity

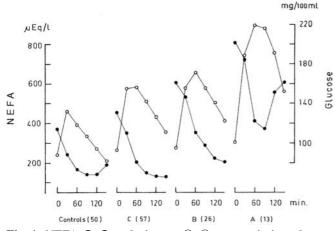
The major change in the appreciation of the nature of molecules supplying body energy came with the discoveries by Dole [1] and Gordon and Cherkes [2] that plasma NEFA are the major lipid transporters in the mammalian organism. Furthermore, NEFA are the only form of fat release from adipose tissue, subject to hormonal regulation [3]. The very rapid turnover of plasma NEFA indicates that they are an important fuel, which even in non-fasting conditions are the energy source to body tissues equivalent to or greater than glucose. Early data on the role of NEFA in glucose homeostasis were reviewed in 1969 by Ruderman et al. [4]. The importance of FFA in the regulation of gluconeogenesis was extensively dealt with, but no clear pathogenic conclusion with relation to major diabetes perturbations emerged at that time.

The effect of hyperinsulinaemia after a glucose load on patterns of NEFA response in normal and diabetic subjects is shown (Fig. 1). The gradual decrease of plasma NEFA concentrations reached a nadir of about 200 µmol/l at 1 h. Non-obese patients with mild Type 2 diabetes showed a decrease in NEFA to a similar and even more protracted nadir than normal subjects most probably because of facilitated uptake of NEFA by tissues in the presence of compensatory hyperinsulinaemia. Patients of two additional groups with high resistance to insulin responded with a more limited NEFA decline pattern and with a higher and shorter NEFA nadir indicating a sustained presence of high NEFA in the circulation and insulin resistance in adipose tissue. The NEFA response is highly sensitive and provides information supplementary to the glucose curve on the patient's response in this situation [5].

Obesity or diabesity as such is another example of fat metabolism abnormality in Type 2 diabetes, whether

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**Fig. 1.** NEFA ●–● and glucose  $\bigcirc$ – $\bigcirc$  curves during glucose load of 1 g/kg in 50 control and 96 patients with IGT or Type 2 diabetes, all non-obese (BMI 24 to 28 kg/m<sup>2</sup>). The patterns of NEFA decline where divided as "Group C" (*n*=57) – nadir similar to controls but occurring later than in control patients, at 60 min and lasting up to 150 min. "Group B" (*n*=26) NEFA decrease not reaching the nadir of the control groups even at 150 min. "Group A" (*n*=13) nadir of NEFA much higher than normal, lasting only 30 min between 60 and 90 min, most probably related to the high preload NEFA value. The higher initial concentrations of NEFA and the high nadirs in groups A and B indicate lower lipolysis restraint in adipose tissue in addition to the impaired glucose tolerance (unpublished investigations of E. Shafrir and A. Gutman based on [5])

due to deficiency of lipostatic hormone or to insulin + glucose promotion of fat deposition. The mechanism of adipose tissue weight gain will not be elaborated here, although diabesity should definitely be looked upon as an aspect of fat metabolism aberration.

It is important to mention at this point that Zierler and Rabinowitz [6] showed in 1964, that infusion of a very small amount of insulin (10 µunits·min<sup>-1</sup>·kg<sup>-1</sup>) through the brachial artery in human forearm leads to a significant decrease in plasma NEFA concentrations and an increase in tissue potassium uptake without any concomitant effect on glucose uptake. This finding leads to the conclusion that the restraint of NEFA release (from adipose tissue) is the most sensitive action of insulin on the lipolysis regulation in comparison with other insulin effects and does not involve the translocation of glucose to promote NEFA retention by esterification.

The lack of restraint of NEFA mobilisation in hypoinsulinaemia leads to a marked plasma NEFA increase, ectopic deposition of triglycerides, insulin resistance and defect in glucose uptake by muscles. The latter fact stresses the effect of excessive availability of NEFA on the reduction of the insulin-mediated muscle glucose utilisation as shown in the classic experiments of Randle et al. [7, 8]. It involves a reduction of muscle glycolysis by inhibiting phosphofructo-kinase by rising cytosolic concentrations of citrate and consequent accumulation of glucose 6-phosphate. Pyruvate dehydrogenase is also inhibited by rising

mitochondrial concentrations of acetylCoA and the NADH:NAD ratio. In the liver, the same changes in redox ratio and activation of pyruvate carboxylase by acetylCoA result in stimulation of gluconeogenesis. This is a further instance of the effect of NEFA oversupply as a contribution to hyperglycaemia in diabetes.

Life-threatening events can occur when the heart NEFA inflow becomes excessive. In acute myocardial ischaemia, it is often associated with fatal ventricular arrhythmia and fibrillation. NEFA have been implicated as a risk factor for sudden cardiac death in diabetes [9, 10]. The arrythmogenic effect of NEFA, related to acute coronary syndrome, is elicited by rapid catecholamine release and massive NEFA mobilisation from adipose tissue. The regional switch in myocardial substrate metabolism to increased fat use elicits reduction in glucose uptake, resistance to insulin action and increase in mitochondrial NEFA oxidation, leading to oxygen crisis.

In pronounced hypoinsulinaemia, when the NEFA supply from adipose tissue escapes control, not only total body glucose homeostasis is disrupted, but the resultant ketosis and acidosis become life-threatening. It is of interest to recall here McGarry's thoughts of Minkowski's failure to identify ketonuria [11] during his seminal experiments with dogs after pancreatectomy [12]. McGarry hypothesised that if Minkowski had been aware of the ketone odour of urine rather than of its sweetness, the importance of the deranged fat metabolism in diabetes would have been recognised sooner.

The eminent Randle glucose-NEFA cycle concept appears, however, to hold only in severe NEFA oversupply in conditions similar to Type 1 diabetes. The effect of lipid infusion in healthy volunteers and the results of a glucose load in obese and insulin-resistant diabetic patients in a euglycaemic clamp system have been investigated [13]. The increased NEFA concentrations in the circulation in these groups, in the presence of insulin, resulted only in relatively moderate decreases in glucose uptake, non-oxidative metabolism and glucose storage with greater utilisation of the fat substrate. The hepatic glucose production was not abolished, which in the presence of increased NEFA availability became non-suppressible by insulin.

The causes for the diminished muscle glucose uptake before the onset of overt diabetes have been extensively reviewed [14, 15, 16]. Convincing evidence that an increase in NEFA concentration, whether due to increased endogenous mobilisation or to postabsorptive lipolysis on a fat-rich diet, is associated with triglyceride deposition in muscle, liver and other tissues, eliciting insulin resistance. Other studies have also indicated that the early perturbations in glucose metabolism as a result of augmented fatty acid utilisation during lipid infusion are gradual [17]. Whole body glucose uptake decreased to about one half of the control after 6 h, augmented lipid oxidation was associated with a fourfold reduction in oxidative glucose metabolism, including glycogen synthesis by 3 h and the decrease in muscle glucose-6-phosphate concentration already by 1.5 h of infusion. Further evidence has shown that NEFA reduce the insulin-stimulated glucose transport and phosphorylation and whole body glucose clearance in human volunteers [18]. Thus, preceding the changes described by Randle et al. [7], the first effect of FFA-induced insulin resistance is the restriction of glucose cell entry, followed by reduction in its metabolism, including the incorporation into glycogen.

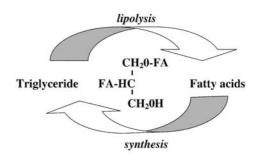
The intracellular muscle triglyceride deposition in subjects susceptible to diabetes was documented by numerous investigators [19, 20, 21, 22, 23] and shown to attenuate glucose metabolism. The earliest observed perturbations are retardation in GLUT4 trafficking reducing the glucose entry into muscle cells [24]. These early changes indicate that phosphatidylinositol-3-kinase (PI3K), regulating GLUT4 translocation to the cell membrane, is affected by reduced IRS tyrosine phosphorylation. Reduction in muscle fat by diet restriction in experimental models of diabesity, the OLETF rats [25] caused an impressive improvement in IGT and in muscle glucose uptake. Similar results were reported in patients with morbid obesity, in which dietary fat deprivation lowered muscle triglycerides, as assessed by histochemistry, ameliorated insulin resistance and increased GLUT4 expression [26]. Also, weight loss after gastric bypass surgery in morbid obese patients resulted in increased insulin sensitivity and reduction in muscle long chain fatty acid CoA (LCFACoA) esters [27].

As discussed in depth by McGarry [28], accumulation of muscle triglycerides results in an increased intracellular LCFACoA concentration, due to lipolysis of the enlarged cell pool of triglycerides. LCFACoA lead to impairment of glucose uptake and insulin resistance through the inhibition of mitochondrial palmitoyl carnitine transferase (CPT) responsible for their transport into the mitochondria and oxidation. The delay in LCFACoA transport causes an increase in the cellular intermediates of triglyceride synthesis and breakdown, with accumulation of intermediates, particularly diacylglycerol (DAG) [29, 30] (Fig. 2). A similar increase in LCFACoA occurs by inhibiting CPT with etomoxir [31]. DAG is not only an outcome of increased triglyceride turnover, but could also be produced by phospholipase C cleavage of diacylinositol phosphates.

### Causative alterations in tissue fat metabolism leading to Type 2 diabetes: DAG and PKC

The outcome of LCFCoA and DAG accumulation is the attenuation of the insulin signalling pathway at

Muscle fat turnover

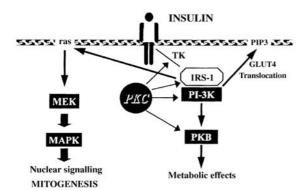


PIP<sub>2</sub> ..... DAG + IP<sub>3</sub>

**Fig. 2.** Source of tissue diacylglycerol (DAG). Synthesis of triglycerides from glucose or intracellular lipolysis of preformed triglycerides involves diacylglycerol as an intermediate. This could also include long-chain fatty acyl CoA (LCFACoA) which accumulate within the cells when their entry into the mitochondria is retarded. Action of phospholipase C on phosphatidylinositol-diphosphate (PIP<sub>2</sub>)

several points. DAG induces an overexpression of isoenzymes of the PKC group, e.g. PKC $\theta$  [32] and PKC $\epsilon$ [33, 34, 35]. Referring to the conjunction of muscle triglycerides, DAG and PKC, in vitro uptake of NEFA by smooth muscle cells increases DAG concentrations with time of incubation and with NEFA concentration [36]. An extensive review of the effects of DAG-sensitive and DAG-independent members of the PKC family is available [37].

The DAG-sensitive PKC isoenzymes are known to phosphorylate serine and threonine on several protein components of the insulin signalling pathway causing a negative feedback in signal transduction [38, 39, 40, 41, 42]. Inhibition of PKC activity was shown to relieve the insulin signalling attenuation [43]. One of the results of PKC-DAG interaction is the reduction of glucose uptake in insulin-sensitive tissues as manifested by inhibition of P13K detachment from the IRS and failure to activate and/or translocate GLUT4 vesicles. A strongly reduced concentration of GLUT4 protein and mRNA was seen in the insulin-resistant PKCe overexpressing *Psammomys obesus* [33, 34], a desert gerbil, prone to nutritionally-induced insulin resistance. The increase in muscle DAG was associated with deficient activation of insulin receptor tyrosine kinase by insulin [44], inhibition of PKB/akt, which regulates numerous metabolic systems within the cell, and degradation of insulin receptor [34, 35]. Apart from PKCE overexpression there was a shift in its cellular distribution towards the membrane, indicating its activation. PKC $\epsilon$  and PKC $\alpha$  were correlated with muscle DAG content in *Psammomys* [35] and PKC0 in high fat fed rats [45, 46]. However, in cultured myotubes obtained from IGT patients, the activation of PI3K and a non-DAG sensitive PKC in response to insulin was decreased [47], which was interpreted as inducing insulin resistance. In human subjects an



**Fig. 3.** Protein kinase C and insulin signaling pathway. The scheme illustrates the association of PKC overexpression (in particular PKCɛ in *Psammomys obesus*) and the attenuating action on insulin receptor tyrosine kinase (TK), phosphatidyl-inositol-3-kinase (PI3K) regulating the translocation of GLUT4 to the membrane and protein kinase B/akt. (Reproduced with permission from [35])

increase in circulating NEFA for a few hours, by lipid + heparin infusion, led to muscle DAG and PKC rise and insulin resistance [32]. The overall effect of PKC on muscle insulin signal transduction is shown in Fig. 3.

Muscle also contains a specific acetyl-CoA carboxylase, which produces malonyl-CoA. However, malonyl-CoA is not used further for fatty acid synthesis as muscles are deficient in fatty acid synthase effecting fatty acid elongation. Malonyl-CoA in muscle is a sensor of NEFA oxidation, its concentrations regulating the entry of LCFACoA into the mitochondria [48, 49]. In turn, the concentrations of malonyl-CoA, apart from acetyl-CoA carboxylase, are also regulated by malonyl-CoA decarboxylase and AMP-activated protein kinase [50]. In muscle, when acetyl-CoA carboxvlase and malonyl-CoA are overexpressed, the entry of NEFA into the mitochondria and their oxidation becomes retarded and LCFACoA rise in concentration. As a result, DAG accumulation ensues, inducing PKC activation and insulin resistance by attenuation of the signalling pathway. Evidence of lower than normal muscle capacity to oxidise fatty acids in the clinical situation is also available. Obese women who reverted to normal weight after caloric restriction, did not respond to a fatty meal with appropriate fat oxidation compared with matched non-obese women [51]. A reduced capacity to oxidise fat was also shown in muscle samples of obese subjects [52]. Despite considerable uptake of palmitate, the muscle of diabetic patients released fewer oxidation products than healthy control subjects [53]. In contrast, muscle fat oxidation was enhanced when rat muscle acetyl-CoA carboxylase, malonyl-CoA decarboxylase and AMP protein kinase responded in opposite direction as a result of exercise [54].

# Other lipid substrates and cytokines involved in attenuation of insulin signaling

It has been shown that the spingolipid ceramide, a major membrane component, can undergo lipolysis by sphigomyelinase, activated by TNF $\alpha$  [55, 56]. This cytokine is secreted by adipose tissue and other tissues in diabesity and effects an inhibition of PKB/akt, IRS and of other components of the insulin signalling pathway in cultured muscle cells [57, 58]. Placental TNF $\alpha$ , mostly released into the maternal circulation, is considered to be partly responsible for the gestational insulin resistance in human pregnancy [59]. Adipose tissue also shows an endocrine-like activity and the abdominal fat is one of the most important risk factors for Type 2 diabetes. In addition to  $TNF\alpha$ , it is a prominent source of cytokines affecting tissue insulin sensitivity and beta-cell function [60]. They involve interleukin-6 [61], resistin [62] and adiponectin [63].

## Diabetes complications: atherosclerosis and hyperlipidaemia

Among the diabetes complications, there is a pronounced acceleration of atherosclerosis, another process based on fat metabolism abnormality. Hyperglycaemia results in a plethora of advanced glycation products, giving rise to reactive oxygen molecules, which activate monocytes and macrophages, causing the proliferation of vascular smooth muscle cells and an overall diabetic angiopathy. One of the main actions in this process is the oxidation of the unsaturated fatty acids esterified to cholesterol in LDL [64]. The oxidized hydroperoxides of cholesterol esters impart a structural change in their apoB protein carrier and thus the LDL bind to surrogate scavenging receptors on macrophages due to nonrecognition by the genuine receptors [65]. The LDL accumulate in atherosclerotic plaques by association with intimal proteoglycans, facilitate macrophage - foam cell conversion, endothelial deposition and smooth muscle cell proliferation [66]. The presence of the scavenger receptor in smooth muscle cells is increased by PKC $\alpha$  [67]. The increase in VLDL and LDL emphasises the role of cholesterol oxidation in the accelerated atherogenesis in diabetes.

Dyslipidaemia plays an important role in diabetic complications, especially cardiovascular disease [68, 69]. The major lipoprotein abnormality in Type 2 diabetes is increased VLDL synthesis, in part due to the hepatic recirculation of NEFA outflowing in excess from adipose tissue, and in part as a result of de novo synthesis in the presence of hyperglycaemia and hyperinsulinaemia. This occurs because the hepatic VLDL synthesis, usually suppressed by insulin, continues unabated in the condition of insulin resistance [70]. There is also an impediment in VLDL and chylomicron removal due to lower than normal lipoprotein lipase (LPL) activity in adipose tissue in diabetes [71, 72, 73]. Changes in VLDL composition, such as excessive triglyceride and low apoE contents [74], can also reduce the LPL activity.

The volume of VLDL particles is of major importance in the atherogenicity in diabetes by determining the size and cholesterol content of the ensuing LDL particles. Large VLDL particles with high triglyceride content are the precursors of small, dense LDL, which determine the severity of atherosclerosis [76, 77]. The activity of cholesterol ester transfer protein (CETP), exchanging triglyceride for cholesterol in hypertriglyceridemic VLDL is increased in diabetes [78]. CETP could contribute to the formation of atherogenic lipoprotein phenotype of diabetes by producing VLDL particles rich in cholesterol and HDL rich in triglyceride not functioning well in the uptake of cholesterol from LDL and from the vessel wall [79].

VLDL in diabetes contain increased Apo B48 of intestinal origin [80, 81], indicating that the diabetic dyslipidaemia is a postprandial metabolic disorder. Improvement in diabetic control reduces the ApoB48 concentrations in VLDL [82]. The avid binding of ApoB48 to the intima, smooth muscle cells and macrophages indicates the presence of high receptor affinity and could explain the preferential appearance of ApoB48 derived from the large VLDL in the atherosclerotic plaques [83, 84].

#### **Diabetic complications: fatty acid** and **PKC-related complications**

Hyperglycaemia in poorly controlled Type 1 diabetes or beta-cell deficient Type 2 diabetes is the main reason for complications as established in the DCCT and UKPDS studies. Nephropathy, neuropathy and retinopathy, have also been indicated as associated with increased oxidative stress and flawed fat metabolism, including vascular contractility, basement membrane thickening, extracellular matrix expansion, microangiopathy and macroangiopathy. Many biochemical and pathological lesions in these insulin independent tissues have been associated with the DAG-sensitive PKC isoenzymes and have been related to changes in Na<sup>+</sup>K<sup>+</sup> ATPase activity, and MAP kinase. Among the PKC isoenzymes overexpressed in these tissues and actively investigated with a specific inhibitor are PKC $\beta_2$  and PKC $\delta$  [85].

### NEFA, triglyceride and beta-cell lipotoxicity

NEFA are known to be insulinotropic [86]. This property represents a very important protection against hypoinsulinaemia on fasting, when the glucose is not available for the stimulation of the pancreatic insulin secretion. The increased mobilisation of NEFA from adipose tissue provides enough insulin to ensure a minimal insulin level necessary for preventing unrestrained NEFA outpour with ketosis.

However, a large inflow of NEFA and an accumulation of triglycerides in beta cells have a negative effect on insulin secretion. In massive obesity with excessive NEFA flow the increased secretion pressure results in beta-cell deterioration up to apoptosis. This has been shown in ZDF rats on relatively high fat diets [87], in *Psammomys obesus* on a high energy diet in the terminal stage characterized by hypoinsulinaemia and an increase in plasma NEFA and triglyceride concentrations [88] and in isolated islets exposed to high NEFA concentrations [89, 90]. The triglyceride content of beta cells increased, intracellular lipolysis raised the NEFA concentration, insulin stores became depleted and the response to glucose was minimal. In human islets preculturing with NEFA caused a caspase-mediated apoptosis possibly involving the ceramide pathway [91].

#### Fat metabolism targeted antidiabetic modalities

Most drugs used in Type 2 diabetes affect fat metabolism albeit in an indirect fashion. However, one of the prominent mechanisms for the increase of insulin sensitivity in muscle and liver by thiazolidinediones (TZD) is their effect on PPAR $\gamma$  receptors in adipose tissue. They promote preadipocyte proliferation, sensitise the inhibition of triglyceride lipolysis by insulin [92, 93, 94], which results in triglyceride retention in adipose tissue and reduction of NEFA availability in the circulation. As discussed above, this prevents the ectopic fat deposition and the development of insulin resistance. Similarly, TZD reduce NEFA oxidation and glucose production in the liver [95]. Perhaps the most impressive model showing this aspect of TZD action is the fat-depleted transgenic mouse model [96, 97]. Because of the non-availability of fat substrate, muscle insulin responsiveness was very high but could be reversed by fat implantation. Similarly, the surgical excision of visceral fat in ageing Zucker diabetic and FBN insulin-resistant rats markedly improved the peripheral and hepatic insulin action [98]. In high fat fed rats, treatment with another TZD, BRL49653, reduced systemic NEFA and pronouncedly improved insulin action and glucoregulation [99]. Treatment with rosiglitazone of Type 2 diabetic patients resulted in a shift of skeletal muscle, mesenteric and hepatic and beta-cell triglyceride content to extracellular sites; hyperinsulinaemic-euglycaemic clamp demonstrated a remarkable improvement in glucose metabolism and an inhibitory activity on adipocyte lipolysis [100].

### Conclusion

We invite comments as to what extent the accumulating evidence indicates that multiple aberrations in fat metabolism, reviewed above, play a pivotal role in the abnormalities of glucose homeostasis in diabetes. In our view the new findings of ectopic fat deposition in muscles and other tissues, linked to DAG accumulation and activation of DAG-sensitive PKC isoenzymes, are responsible for the emerging insulin resistance and beta-cell dysfunction, with the transition to full-fledged Type 2 diabetes. These findings implicate the derangements in fat metabolism as the main culprit of metabolic deviations in Type 2 diabetes: the negative feedback of the insulin signalling pathway. We maintain that these findings clarify the pathophysiology of diabetes and require redefinition of the glucocentric versus lipocentric approach. Our recommendation is as a first step to drop the adjective "mellitus" from diabetes and then to consider the introduction of a new adjective "lipidus" or "lipomellitus".

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